

## SEQUENCE LISTING

<110> Johnson, Jason M.  
Castle, John C.  
Armour, Christopher D.

<120> SPLICE VARIANT ISOFORMS OF HUMAN CALCIUM CHANNEL CACNA1B

<130> RS0202Y

<160> 13

<170> PatentIn version 3.2

<210> 1  
<211> 6792  
<212> DNA  
<213> Homo sapiens

<400> 1  
atggccgct tcggggacga gctgggcggc cgctatggag gccccggcgg cggagagcgg 60  
gccccgggacg gcggggccgg cggggcgggg ggcccgggtc ccggggggct gcagccggc 120  
cagcgggtcc tctacaagca atcgatcgcg cagcgcgcgc ggaccatggc gctgtacaac 180  
cccatcccg tcaagcagaa ctgcttcacc gtcaaccgct cgctttcgt cttcagcgg 240  
gacaacgtcg tccgcaaata cgcgaagcgc atcaccgagt ggccctccatt cgagtatatg 300  
atcctggcca ccatcatcgc caactgcattc gtgctggccc tggagcagca cctccctgat 360  
ggggacaaaa cgcgcattgtc cgagcggctg gacgacacgg agccctattt catcgggatc 420  
ttttgcttcg aggcaaggat caaaatcatc gctctggct ttgtcttcca caagggctct 480  
tacctgcgga acggctggaa cgtcatggac ttctgtggcg tcctcacagg gatccttgc 540  
acggctggaa ctgacttcga cctgcgaaca ctgagggctg tgcgtgtgct gaggccccctg 600  
aagctgggt ctgggattcc aagttgcag gtggtgctca agtccatcat gaaggccatg 660  
gttccactcc tgcagattgg gctgtttctc ttctttgcca tcctcatgtt tgccatcatt 720  
ggcctggagt tctacatggg caagttccac aaggcctgtt tccccaaacag cacagatgcg 780  
gagccccgtgg gtgacttccc ctgtggcaag gaggccccag cccggctgtg cgagggcgcac 840  
actgagtgcc gggagttactg gccaggaccc aactttggca tcaccaactt tgacaatatc 900  
ctgtttgcca tcttgacggt gttccagtgc atcaccatgg agggctggac tgacatcctc 960  
tataatacaa acgatgcggc cggcaacacc tggaaactggc tctacttcattt ccctctcatc 1020  
atcatcggt ctttcttcattt gctcaacccgtt gtgctggcg tgctctcggg ggagtttgc 1080  
aaggagcgag agaggggtgga gaaccgcgc gccttcgttga agctgcgcgg gcagcagcag 1140  
atcgagcgag agctcaacgg gtacctggag tggatcttca aggcggagga agtcatgtg 1200  
gccgaggagg acaggaatgc agaggagaag tcccccttgg acgtgctgaa gagagcggcc 1260

accaagaaga	gcagaaaatga	cctgatccac	gcagaggagg	gagaggaccg	gtttgcagat	1320
ctctgtgctg	ttggatcccc	cttcgcccgc	gccagcctca	agagcgggaa	gacagagagc	1380
tcgtcatact	tccggaggaa	ggagaagatg	ttccgggttt	ttatccggcg	catggtgaag	1440
gctcagagct	tctactgggt	ggtgctgtgc	gtgggtggccc	tgaacacact	gtgtgtggcc	1500
atggtgcatt	acaaccagcc	gcggcggctt	accacgaccc	tgtatttgc	agagtttgtt	1560
ttcctgggtc	tcttcctcac	agagatgtcc	ctgaagatgt	atggcctggg	gcccagaagc	1620
tacttccgg	cctccttcaa	ctgcttcgac	tttggggtca	tcgtggggag	cgtctttgaa	1680
gtggtctggg	cggccatcaa	gccgggaagc	tcctttggga	tcagtgtgct	gcgggcccctc	1740
cgcctgctga	ggatcttcaa	agtcacgaag	tactggagct	ccctgcggaa	cctgggtgg	1800
tccctgctga	actccatgaa	gtccatcatc	agcctgctct	tcttgctctt	cctgttcatt	1860
gtggtcttcg	ccctgctggg	gatgcagctg	tttgggggac	agttcaactt	ccaggatgag	1920
actcccacaa	ccaaacttcga	cacctccct	gccgccatcc	tcactgtctt	ccagatcctg	1980
acgggagagg	actggaatgc	agtgatgtat	cacgggatcg	aatcgcaagg	cggcgtcagc	2040
aaaggcatgt	tctcgtcctt	ttacttcatt	gtcctgacac	tgttcggaaa	ctacactctg	2100
ctgaatgtct	ttctggccat	cgctgtggac	aacctggcca	acgcccaga	gctgaccaag	2160
gatgaagagg	agatggaaga	agcagccaat	cagaagctt	ctctgcaaaa	ggccaaagaa	2220
gtggctgaag	tcaaaaaat	gtctgccgcg	aacatctcca	tcgcccag	gcagcagaac	2280
tcggccaagg	cgcgctcggt	gtgggagcag	cggggcagcc	agctacggct	gcagaacctg	2340
cggggcagct	gcgaggcgct	gtacagcgag	atggaccccg	aggagcggct	gcgttcgccc	2400
actacgcgcc	acctgcggcc	cgacatgaag	acgcacctgg	accggccgct	ggtgggtggag	2460
ctggccgcgc	acggcgcgcg	ggggcccgtg	ggaggcaaag	cccgacctga	ggctgcggag	2520
gcccccgagg	gcgtcgaccc	tccgcgcagg	caccacccgc	accgcgacaa	ggacaagacc	2580
cccgccgggg	gggaccagga	ccgagcagag	gccccgaagg	cggagagcgg	ggagcccggt	2640
gccccggagg	agcggcccg	gccgcaccgc	agccacagca	aggagccgc	ggggcccccg	2700
gaggcgcgga	gcgagcgcgg	ccgaggccca	ggccccgagg	gcggccggcg	gcaccacccgg	2760
cgcgcgtccc	cggaggagggc	ggccgagcgg	gagccccgac	gccaccgcgc	gcaccggcac	2820
caggatccga	gcaaggagtg	cgccggcgcc	aagggcgagc	ggcgcgcgcg	gcaccgcggc	2880
ggcccccggag	cggggcccccg	ggaggcggag	agcggggagg	agccggcg	gcggcaccgg	2940
gccccgcaca	aggcgcagcc	tgctcacgag	gctgtggaga	aggagaccac	ggagaaggag	3000
gccacggaga	aggaggctga	gatagtggaa	gccgacaagg	aaaaggagct	ccggaaccac	3060

cagccccggg agccacactg tgacctggag accagtggga ctgtgactgt	3120
cacacactgc ccagcacctg tctccagaag gtggaggaac agccagagga	3180
cagcggAACG tcactcgcat gggcagtcag ccccccagacc cgaacactat	3240
ccagtgatgc tgacggggcc tcttgggaa gccacggtcg ttcccagtgg	3300
ctggaaagcc aagcagaggg gaagaaggag gtggaaagcgg atgacgtgat	3360
ccccggccta tcgtccccata cagctccatg ttctgtttaa gccccaccaa	3420
cgttctgcc actacatcg gaccatgagg tacttcgagg tggtcattct	3480
gccttgagca gcatcgccct ggctgctgag gacccagtgc gcacagactc	3540
aacgctctga aatacctgga ttacatttc actgggtgtct ttaccttga	3600
aagatgatcg acttgggact gctgcttcac cctggagcct atttccggga	3660
attctggact tcattgtggt cagtggcgcc ctgggtggcgt ttgcttctc	3720
gggaaagaca tcaataccat caagtctctg agagtccctc gtgtcctgcg	3780
accatcaaac ggctgccccaa gctcaaggct gtgtttgact gtgtggtcaa	3840
aatgtcctca acatcttcatg tgtctacatg ctcttcatgt tcataattgc	3900
gtgcagctct tcaaaggaa gttttctac tgacacagatg aatccaagga	3960
gactgcaggg gtcagtttggatttggatg aaggaggaag tggaagctca	4020
gtggaaat acgactttca ctacgacaat gtgctctggg ctctgctgac	4080
gtgtccacgg gagaaggctg gcccattggtg ctgaaacact ccgtggatgc	4140
gagcagggtc caagccctgg gtaccgcattg gagctgtcca tcttctacgt	4200
gtggctttc cttttttttt cgtcaacatc tttgtggctt tgatcatcat	4260
gagcaggggg acaagggtat gtctgaatgc agcctggaga agaacgagag	4320
gacttcgcaca tcagcgccaa accccatgaca cggtacatgc cccaaaaccc	4380
cagtataaga cgtggacatt tgggtctcc ccgcctttt aataacttcat	4440
atagccctca acactgtggt gctgatgatg aagttctatg atgcacccta	4500
ctgatgctga aatgcctgaa catcggttc acatccatgt tctccatgga	4560
aagatcatcg ctttgggggt gctgaactat ttcaagatg cctgaaatgt	4620
gtcactgtgt tggaaagtat tactgatatt ttagtaacag agattgcgg	4680
ttcatcaacc tcagcttcct ccgcctttt cgagctgcgc ggctgatcaa	4740
cagggctaca ccatccgcat cctgctgtgg acctttgtcc agtcctcaa	4800
tacgtgtgtc tgctcattgc catgctgttc ttcatctacg ccatcatcg	4860
tttggaaata ttgcctgga tgatgacacc agcatcaacc gccacaacaa	4920

ttttgcaag ccctgatgct gctgttcagg agcgccacgg gggaggcctg gcacgagatc	4980
atgctgtcct gcctgagcaa ccaggcctgt gatgagcagg ccaatgccac cgagtgtgga	5040
agtactttg cctacttcta cttcgctcctt ttcatcttcc tgtgctcctt tctgcgcctg	5100
gttcgcacatga acatgccat ctccaaacgag gacatgactg ttcaacttcac gtccacgctg	5160
atggccctca tccggacggc actggagatc aagctggccc cagctggac aaagcagcat	5220
cagtgacg cggagtttag gaaggagatt tccgttgtgt gggccaatct gccccagaag	5280
actttggact tgctggtacc accccataag cctgatgaga tgacagtggg gaaggtttat	5340
gcagctctga tgatatttga cttctacaag cagaacaaaa ccaccagaga ccagatgcag	5400
caggctcctg gaggcctctc ccagatgggt cctgtgtccc tttccacccc tctgaaggcc	5460
accctggagc agacacagcc ggctgtgctc cgaggagccc gggttttcct tcgacagaag	5520
agttccaccc ccctcagcaa tggcggggcc atacaaaacc aagagagtgg catcaaagag	5580
tctgtctcct ggggcactca aaggacccag gatgcaccccc atgaggccag gccacccctg	5640
gagcgtggcc actccacaga gatccctgtg gggcggtcag gagcactggc tgtggacgtt	5700
cagatgcaga gcataacccg gaggggccct gatggggagc cccagctgg gctggagagc	5760
cagggtcag cggcctccat gccccccctt gggccgaga ctcagccctg cacagatgcc	5820
agccccatga agcgctccat ctccacgctg gcccagcggc cccgtggac tcacatttgc	5880
agcaccaccc cggaccgccc acccccattgc caggcgtcgt cgcaccacca ccaccaccgc	5940
tgccaccgccc gcagggacag gaagcagagg tccctggaga agggggccag cctgtctgcc	6000
gatatggatg gcgaccaag cagtgtgtg gggccggggc tgccccccgg agaggggcct	6060
acaggctgcc ggcgggaacg agagcgccgg caggagcggg gccgggtccca ggagcggagg	6120
cagccctcat cctcctcctc ggagaagcag cgcttctact cctgcgaccc ctttggggc	6180
cgtgagcccc cgaagcccaa gcccctccctc agcagccacc caacgtcgcc aacagctggc	6240
caggagccgg gaccccaccc acagggcagt gttccgtga atgggagccc cttgctgtca	6300
acatctggtg cttagcaccctt cggccgcggt gggcgaggc agctccccca gacccctcg	6360
actccccgcc ccagcatcac ctacaagacg gccaactcct cacccatcca cttcgccggg	6420
gctcagacca gcctccctgc cttctccca gggcggtca gccgtgggt ttccgaacac	6480
aacgcccctgc tgcagagaga cccctcagc cagccccctgg cccctggctc tcgaattggc	6540
tctgaccctt acctggggca gcgctggac agtgaggcct ctgtccacgc cctgcctgag	6600
gacacgctca ctttcgagga ggctgtggcc accaactcgg gccgctcctc caggacttcc	6660
tacgtgtcct ccctgacccctc ccagtcac cctctccggcc gcgtgcccgg cggttaccac	6720

tgacccctgg gactcagctc gggtgccga gcacggcaca gctaccacca ccctgaccaa 6780  
 gaccactggt gc 6792

<210> 2  
 <211> 2264  
 <212> PRT  
 <213> Homo sapiens

<400> 2

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly  
 1 5 10 15

Gly Gly Glu Arg Ala Arg Gly Gly Ala Gly Gly Ala Gly Gly Pro  
 20 25 30

Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser  
 35 40 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val  
 50 55 60

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu  
 65 70 75 80

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro  
 85 90 95

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu  
 100 105 110

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu  
 115 120 125

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu  
 130 135 140

Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser  
 145 150 155 160

Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr  
 165 170 175

Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg  
 180 185 190

Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser  
 195 200 205  
 Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu  
 210 215 220  
 Gln Ile Gly Leu Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile  
 225 230 235 240  
 Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn  
 245 250 255  
 Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala  
 260 265 270  
 Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Pro  
 275 280 285  
 Gly Pro Asn Phe Gly Ile Thr Asn Phe Asn Ile Leu Phe Ala Ile  
 290 295 300  
 Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu  
 305 310 315 320  
 Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe  
 325 330 335  
 Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu  
 340 345 350  
 Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn  
 355 360 365  
 Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Ile Glu Arg Glu  
 370 375 380  
 Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu  
 385 390 395 400  
 Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu  
 405 410 415  
 Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu  
 420 425 430

Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe  
435 440 445

Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe  
450 455 460

Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys  
465 470 475 480

Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr  
485 490 495

Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Arg Arg Leu Thr Thr  
500 505 510

Thr Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu  
515 520 525

Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser  
530 535 540

Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Val Phe Glu  
545 550 555 560

Val Val Trp Ala Ala Ile Lys Pro Gly Ser Ser Phe Gly Ile Ser Val  
565 570 575

Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp  
580 585 590

Ser Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser  
595 600 605

Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala  
610 615 620

Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu  
625 630 635 640

Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val  
645 650 655

Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly  
660 665 670

Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr  
 675 680 685  
 Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe  
 690 695 700  
 Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys  
 705 710 715 720  
 Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln  
 725 730 735  
 Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile  
 740 745 750  
 Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp  
 755 760 765  
 Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys  
 770 775 780  
 Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala  
 785 790 795 800  
 Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro  
 805 810 815  
 Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly  
 820 825 830  
 Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro  
 835 840 845  
 Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly  
 850 855 860  
 Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly  
 865 870 875 880  
 Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala  
 885 890 895  
 Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro  
 900 905 910

Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala  
 915 920 925  
 Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser  
 930 935 940  
 Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly  
 945 950 955 960  
 Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala  
 965 970 975  
 Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val  
 980 985 990  
 Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile  
 995 1000 1005  
 Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg  
 1010 1015 1020  
 Glu Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly  
 1025 1030 1035  
 Pro Met His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu  
 1040 1045 1050  
 Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly  
 1055 1060 1065  
 Ser Gln Pro Pro Asp Pro Asn Thr Ile Val His Ile Pro Val Met  
 1070 1075 1080  
 Leu Thr Gly Pro Leu Gly Glu Ala Thr Val Val Pro Ser Gly Asn  
 1085 1090 1095  
 Val Asp Leu Glu Ser Gln Ala Glu Gly Lys Lys Glu Val Glu Ala  
 1100 1105 1110  
 Asp Asp Val Met Arg Ser Gly Pro Arg Pro Ile Val Pro Tyr Ser  
 1115 1120 1125  
 Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys  
 1130 1135 1140

His Tyr Ile Val Thr Met Arg Tyr Phe Glu Val Val Ile Leu Val  
1145 1150 1155

Val Ile Ala Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val  
1160 1165 1170

Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys Tyr Leu Asp Tyr  
1175 1180 1185

Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile  
1190 1195 1200

Asp Leu Gly Leu Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu  
1205 1210 1215

Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala  
1220 1225 1230

Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys  
1235 1240 1245

Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys  
1250 1255 1260

Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser  
1265 1270 1275

Leu Lys Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met  
1280 1285 1290

Phe Ile Phe Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe  
1295 1300 1305

Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg  
1310 1315 1320

Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro  
1325 1330 1335

Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr Asp Asn Val Leu Trp  
1340 1345 1350

Ala Leu Leu Thr Leu Phe Thr Val Ser Thr Gly Glu Gly Trp Pro  
1355 1360 1365

Met Val Leu Lys His Ser Val Asp Ala Thr Tyr Glu Glu Gln Gly  
 1370 1375 1380  
 Pro Ser Pro Gly Tyr Arg Met Glu Leu Ser Ile Phe Tyr Val Val  
 1385 1390 1395  
 Tyr Phe Val Val Phe Pro Phe Phe Val Asn Ile Phe Val Ala  
 1400 1405 1410  
 Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp Lys Val Met Ser  
 1415 1420 1425  
 Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp Phe Ala  
 1430 1435 1440  
 Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn Arg Gln  
 1445 1450 1455  
 Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro Phe  
 1460 1465 1470  
 Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu  
 1475 1480 1485  
 Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu  
 1490 1495 1500  
 Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys  
 1505 1510 1515  
 Val Leu Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp  
 1520 1525 1530  
 Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr  
 1535 1540 1545  
 Asp Ile Leu Val Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn  
 1550 1555 1560  
 Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu  
 1565 1570 1575  
 Leu Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val  
 1580 1585 1590

Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu Ile Ala Met  
1595 1600 1605  
Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val Phe Gly Asn  
1610 1615 1620  
Ile Ala Leu Asp Asp Asp Thr Ser Ile Asn Arg His Asn Asn Phe  
1625 1630 1635  
Arg Thr Phe Leu Gln Ala Leu Met Leu Leu Phe Arg Ser Ala Thr  
1640 1645 1650  
Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys Leu Ser Asn Gln  
1655 1660 1665  
Ala Cys Asp Glu Gln Ala Asn Ala Thr Glu Cys Gly Ser Asp Phe  
1670 1675 1680  
Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser Phe Leu  
1685 1690 1695  
Arg Leu Val Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr  
1700 1705 1710  
Val His Phe Thr Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu  
1715 1720 1725  
Glu Ile Lys Leu Ala Pro Ala Gly Thr Lys Gln His Gln Cys Asp  
1730 1735 1740  
Ala Glu Leu Arg Lys Glu Ile Ser Val Val Trp Ala Asn Leu Pro  
1745 1750 1755  
Gln Lys Thr Leu ASP Leu Leu Val Pro Pro His Lys Pro ASP Glu  
1760 1765 1770  
Met Thr Val Gly Lys Val Tyr Ala Ala Leu Met Ile Phe Asp Phe  
1775 1780 1785  
Tyr Lys Gln Asn Lys Thr Thr Arg Asp Gln Met Gln Gln Ala Pro  
1790 1795 1800  
Gly Gly Leu Ser Gln Met Gly Pro Val Ser Leu Phe His Pro Leu  
1805 1810 1815

Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Val Leu Arg Gly Ala  
 1820 1825 1830  
  
 Arg Val Phe Leu Arg Gln Lys Ser Ser Thr Ser Leu Ser Asn Gly  
 1835 1840 1845  
  
 Gly Ala Ile Gln Asn Gln Glu Ser Gly Ile Lys Glu Ser Val Ser  
 1850 1855 1860  
  
 Trp Gly Thr Gln Arg Thr Gln Asp Ala Pro His Glu Ala Arg Pro  
 1865 1870 1875  
  
 Pro Leu Glu Arg Gly His Ser Thr Glu Ile Pro Val Gly Arg Ser  
 1880 1885 1890  
  
 Gly Ala Leu Ala Val Asp Val Gln Met Gln Ser Ile Thr Arg Arg  
 1895 1900 1905  
  
 Gly Pro Asp Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg  
 1910 1915 1920  
  
 Ala Ala Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Val Thr  
 1925 1930 1935  
  
 Asp Ala Ser Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Gln Arg  
 1940 1945 1950  
  
 Pro Arg Gly Thr His Leu Cys Ser Thr Thr Pro Asp Arg Pro Pro  
 1955 1960 1965  
  
 Pro Ser Gln Ala Ser Ser His His His His His Arg Cys His Arg  
 1970 1975 1980  
  
 Arg Arg Asp Arg Lys Gln Arg Ser Leu Glu Lys Gly Pro Ser Leu  
 1985 1990 1995  
  
 Ser Ala Asp Met Asp Gly Ala Pro Ser Ser Ala Val Gly Pro Gly  
 2000 2005 2010  
  
 Leu Pro Pro Gly Glu Gly Pro Thr Gly Cys Arg Arg Glu Arg Glu  
 2015 2020 2025  
  
 Arg Arg Gln Glu Arg Gly Arg Ser Gln Glu Arg Arg Gln Pro Ser  
 2030 2035 2040

Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr Ser Cys Asp Arg Phe  
 2045 2050 2055  
 Gly Gly Arg Glu Pro Pro Lys Pro Lys Pro Ser Leu Ser Ser His  
 2060 2065 2070 2075  
 Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro His Pro Gln  
 2075 2080 2085  
 Gly Ser Gly Ser Val Asn Gly Ser Pro Leu Leu Ser Thr Ser Gly  
 2090 2095 2100  
 Ala Ser Thr Pro Gly Arg Gly Gly Arg Arg Gln Leu Pro Gln Thr  
 2105 2110 2115  
 Pro Leu Thr Pro Arg Pro Ser Ile Thr Tyr Lys Thr Ala Asn Ser  
 2120 2125 2130  
 Ser Pro Ile His Phe Ala Gly Ala Gln Thr Ser Leu Pro Ala Phe  
 2135 2140 2145  
 Ser Pro Gly Arg Leu Ser Arg Gly Leu Ser Glu His Asn Ala Leu  
 2150 2155 2160  
 Leu Gln Arg Asp Pro Leu Ser Gln Pro Leu Ala Pro Gly Ser Arg  
 2165 2170 2175  
 Ile Gly Ser Asp Pro Tyr Leu Gly Gln Arg Leu Asp Ser Glu Ala  
 2180 2185 2190  
 Ser Val His Ala Leu Pro Glu Asp Thr Leu Thr Phe Glu Glu Ala  
 2195 2200 2205  
 Val Ala Thr Asn Ser Gly Arg Ser Ser Arg Thr Ser Tyr Val Ser  
 2210 2215 2220  
 Ser Leu Thr Ser Gln Ser His Pro Leu Arg Arg Val Pro Asn Gly  
 2225 2230 2235  
 Tyr His Cys Thr Leu Gly Leu Ser Ser Gly Gly Arg Ala Arg His  
 2240 2245 2250  
 Ser Tyr His His Pro Asp Gln Asp His Trp Cys  
 2255 2260

<210>	3					
<211>	5235					
<212>	DNA					
<213>	Homo sapiens					
<400>	3					
atggtccgct	tcggggacga	gctgggcggc	cgctatggag	gccccggcgg	cggagagcgg	60
gccccggg	cgccccgg	cgggggcggg	ggccccgg	ccgggggg	gcagcccc	120
cagcgggtcc	tctacaagca	atcgatcg	cagcgcgc	ggaccatgg	gctgtacaac	180
cccatcccg	tcaagcagaa	ctgcttcacc	gtcaaccg	cgctttcg	cttcagcgag	240
gacaacgtcg	tccgcaaata	cgcgaagcgc	atcaccg	ggcctccatt	cgagtatatg	300
atcctggcca	ccatcatcg	caactgc	gtgctgg	tgagcagca	cctccctgat	360
ggggacaaaa	cgcgcgtc	cgagcgg	gacgacacgg	agccctattt	catcgggatc	420
ttttgttcg	aggcaggat	caaaatc	gtctgg	ttgtcttcca	caagggctct	480
tacctgcg	acggctggaa	cgtcatgg	ttcgtgg	tcctcacagg	gatccttgc	540
acggctggaa	ctgacttcg	cctgc	ctgagg	tgcgtgt	gaggccc	600
aagctgg	ctggattcc	aagttgc	gtgg	gtca	agtccatcat	660
gttccactcc	tgcagattgg	gctgtt	ttcttgc	tcctcatgtt	tgccatcatt	720
ggcctgg	gtctacatgg	caagttcc	aaggc	tttgc	tcccaacag	780
gagccgtgg	gtgacttccc	ctgtgg	gaggccc	ccggctgt	cgagggcg	840
actgagtgc	gggagactg	gccaggac	aacttgg	tcaccaactt	tgacaatatc	900
ctgtttgc	tcttgacgg	gttcc	atcaccatgg	agggctgg	tgacatc	960
tataataca	acgatgc	cgcaacacc	tgaactgg	tctacttcat	ccctctcat	1020
atcatcg	ccttctcat	gctcaac	gtgctgg	tgctctc	ggagtttgc	1080
aaggagc	agaggg	gaaccgc	gccttctg	agctgc	gcagcagc	1140
atcgagc	agctcaac	gtacctgg	tggatcttca	aggcgg	agtcatg	1200
gccgagg	acaggaatgc	agaggaga	tccc	acgtg	gagagcgg	1260
accaaga	gcagaaatga	cctgatcc	gcagagg	gagaggac	gttgc	1320
ctctgt	ttggatcccc	cttcgccc	gccagc	agagcgg	gacagag	1380
tcgtcata	tccggagg	ggagaagat	ttccgg	ttatccgg	catgg	1440
gctcaga	gtctactgg	ggtgctgt	gtgg	tgaacaca	gtgtgtgg	1500
atggtg	cattacaacc	gcggcgg	accacgac	tgtat	ttttgtt	1560
ttcctgg	tcttc	agagatgt	ctgaagatgt	atggc	tgg	1620
tacttc	cctc	ctgcttc	ac	tttgg	cg	1680
gtc	tttgc	ac	tttgc	tttgc	tttgc	

gtggtctggg cggccatcaa gccgggaagc tccttggga tcagtgtgct gcgggccc	1740
ccctgctga ggatcttcaa agtcacgaag tactggagct ccctgcggaa cctgggtgg	1800
tccctgctga actccatgaa gtccatcatc agcctgctct tcttgctctt cctgttcatt	1860
gtggtctcg ccctgctggg gatgcagctg tttggggac agttcaactt ccaggatgag	1920
actcccacaa ccaacttcga cacctccct gccgcatcc tcactgtctt ccagatcctg	1980
acgggagagg actggaatgc agtgatgtat cacggatcg aatcgcaagg cggcgtcagc	2040
aaaggcatgt tctcgccctt ttacttcatt gtcctgacac tgttcggaaa ctacactctg	2100
ctgaatgtct ttctggccat cgctgtggac aacctggcca acgcccaga gctgaccaag	2160
gatgaagagg agatggaaga agcagccaat cagaagctt ctctgcaaaa ggccaaagaa	2220
gtggctgaag tcagccccat gtctgccgcg aacatctcca tcgcccagc gcagcagaac	2280
tcggccaagg cgcgctcggt gtgggagcag cggccagcc agctacggct gcagaacctg	2340
cggccagct gcgaggcgcgt gtacagcag atggaccccg aggagcggct gcgttcgccc	2400
actacgcgcc acctgcggcc cgacatgaag acgcacctgg accggccgct ggtgggtggag	2460
ctggccgcg acggcgccgcg gggcccggt ggaggcaaag cccgacctga ggctgcggag	2520
gcccccgagg gcgtcgaccc tccgcgcagg caccaccggc accgcgacaa ggacaagacc	2580
cccgccggcgg gggaccagga ccgagcagag gccccgaagg cggagagcgg ggagcccggt	2640
gccccggagg agcggccgcg gccgcaccgc agccacagca aggaggccgc gggcccccg	2700
gaggcgcgga gcgagcgcgg ccgaggccca ggccccgagg gcggccggcg gcaccaccgg	2760
cgcggctccc cggaggaggc ggccgagcgg gagcccccgc acgcaccgc gcaccggcac	2820
caggatccga gcaaggagtg cgccggcgcc aaggcgagc ggcgccgcg gcaccgcgc	2880
ggcccccgag cggggccccc ggaggcgag agcggggagg agccggcgcg gcggcaccgg	2940
gccccggcaca aggcgccagcc tgctcacgag gctgtggaga aggagaccac ggagaaggag	3000
gccacggaga aggaggctga gatagtggaa gccgacaagg aaaaggagct ccggaaccac	3060
cagccccggg agccacactg tgacctggag accagtggga ctgtgactgt gggccatg	3120
cacacactgc ccagcacctg tctccagaag gtggaggaac agccagagga tgcagacaat	3180
cagcggAACG tcactcgcat gggcagtcag cccccagacc cgaacactat tgtacatatc	3240
ccagtgtatgc tgacggccccc tcttgggaa gccacggcg ttcccagtgg taacgtggac	3300
ctggaaagcc aagcagaggg gaagaaggag gtggaaagcgg atgacgtat gaggagcggc	3360
ccccggccta tcgtccata cagctccatg ttctgtttaa gccccaccaa cctgctccgc	3420
cgcttctgcc actacatcgat gaccatgagg tacttcgagg tggtcattct cgtggtcatc	3480

gcctttagca gcatcgccct ggctgctgag gacccagtgc gcacagactc gcccaggaac	3540
aacgctctga aatacctgga ttacatttc actgggtctt ttaccttga gatgggtata	3600
aagatgatcg acttgggact gctgcttcac cctggagcct atttccggga cttgtgaaac	3660
attctggact tcattgtggt cagtggcgcc ctgggtggcgt ttgcttctc aggatccaaa	3720
gggaaagaca tcaataccat caagtctctg agagtccttc gtgtctgtcg gcccctcaag	3780
accatcaaac ggctgccc aa gctcaaggct gtgtttgact gtgtggtaa ctccctgaag	3840
aatgtcctca acatcttgcat tgtctacatg ctcttcatgt tcataatttc cgtcattgcg	3900
gtgcagctct tcaaaggaa gttttctac tgcacagatg aatccaagga gctggagagg	3960
gactgcaggg gtcagtattt ggattatgag aaggaggaag tggaaagctca gcccaggcag	4020
tggaaagaaat acgactttca ctacgacaat gtgctctggg ctctgctgac gctgttcaca	4080
gtgtccacgg gagaaggctg gcccattggc ctgaaacact ccgtggatgc caccatgag	4140
gagcagggtc caagccctgg gtaccgcattt gagctgtcca tcttcatactt ggtctacttt	4200
gtggtcttcc cttttttttt cgtcaacatc ttgtggctt tgatcatcat cacccatcag	4260
gagcaggggg acaaggtgat gtctgaatgc agcctggaga agaacgagag ggcttgcatt	4320
gacttcgcca tcagcgccaa acccctgaca cggtacatgc cccaaaaccg gcagtcgttc	4380
cagtataaga cgtggacatt tgtggctcc ccgcctttt aataacttcat catggccatg	4440
atagccctca acactgtggt gctgatgatg aagttctatg atgcacccta tgagtacgag	4500
ctgatgctga aatgcctgaa catcggttc acatccatgt tctccatgaa atcggtgtc	4560
aagatcatcg ctttggggt gctgaactat ttccagatg cctggaaatgt ctttacttt	4620
gtcactgtgt tggaaagttt tactgatatt tttagtaacag agattgcgga aacgaacaat	4680
ttcatcaacc tcagcttcctt ccgcctttt cgagctgcgc ggctgatcaa gctgctccgc	4740
cagggttaca ccatccgtt cctgtgtgg acctttgtcc agtccatcaa ggccctgccc	4800
tacgtgtgtc tgctcattgc catgctgttc ttcatctacg ccatcatcg catgcagggt	4860
tttggaaata ttgcctgga tgatgacacc agcatcaacc gccacaacaa ctccggacg	4920
tttttgcag ccctgatgct gctgttcagg agcgccacgg gggaggcctg gcacgagatc	4980
atgctgtcct gcctgagcaa ccaggctgt gatgagcagg ccaatgccac cgagtgtgga	5040
agtgacttttgcctacttctt cttcgctcc ttcatcttcc tggatgtttt tctgtatgttgc	5100
aaccttttgc tggatgttgcatgatcatggacaat ttgtggatcc tcacgcggga ctccatc	5160
ctaggtcctc accacttggaa tgagttcatc cgggtctggg ctgaatacga cccggctgcg	5220
tgccctgggt tcgca	5235

<210> 4  
<211> 1745  
<212> PRT  
<213> Homo sapiens

<400> 4

Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly  
1 5 10 15

Gly Gly Glu Arg Ala Arg Gly Gly Ala Gly Gly Ala Gly Gly Pro  
20 25 30

Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser  
35 40 45

Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val  
50 55 60

Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu  
65 70 75 80

Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro  
85 90 95

Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu  
100 105 110

Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu  
115 120 125

Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu  
130 135 140

Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser  
145 150 155 160

Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr  
165 170 175

Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg  
180 185 190

Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser  
195 200 205

Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu  
210 215 220

Gln Ile Gly Leu Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile  
225 230 235 240

Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn  
245 250 255

Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala  
260 265 270

Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Pro  
275 280 285

Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile  
290 295 300

Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu  
305 310 315 320

Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe  
325 330 335

Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu  
340 345 350

Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn  
355 360 365

Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu  
370 375 380

Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu  
385 390 395 400

Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu  
405 410 415

Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu  
420 425 430

Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe  
435 440 445

Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe  
450 455 460

Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys  
465 470 475 480

Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr  
485 490 495

Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Arg Arg Leu Thr Thr  
500 505 510

Thr Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu  
515 520 525

Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser  
530 535 540

Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Val Phe Glu  
545 550 555 560

Val Val Trp Ala Ala Ile Lys Pro Gly Ser Ser Phe Gly Ile Ser Val  
565 570 575

Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp  
580 585 590

Ser Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser  
595 600 605

Ile Ile Ser Leu Leu Phe Leu Leu Phe Ile Val Val Phe Ala  
610 615 620

Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu  
625 630 635 640

Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val  
645 650 655

Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly  
660 665 670

Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr  
675 680 685

Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe  
690 695 700

Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys  
705 710 715 720

Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln  
725 730 735

Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile  
740 745 750

Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp  
755 760 765

Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys  
770 775 780

Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala  
785 790 795 800

Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro  
805 810 815

Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly  
820 825 830

Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro  
835 840 845

Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly  
850 855 860

Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly  
865 870 875 880

Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala  
885 890 895

Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro  
900 905 910

Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala  
915 920 925

Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser  
930 935 940

Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly  
945 950 955 960

Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala  
965 970 975

Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val  
980 985 990

Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile  
995 1000 1005

Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg  
1010 1015 1020

Glu Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly  
1025 1030 1035

Pro Met His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu  
1040 1045 1050

Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly  
1055 1060 1065

Ser Gln Pro Pro Asp Pro Asn Thr Ile Val His Ile Pro Val Met  
1070 1075 1080

Leu Thr Gly Pro Leu Gly Glu Ala Thr Val Val Pro Ser Gly Asn  
1085 1090 1095

Val Asp Leu Glu Ser Gln Ala Glu Gly Lys Lys Glu Val Glu Ala  
1100 1105 1110

Asp Asp Val Met Arg Ser Gly Pro Arg Pro Ile Val Pro Tyr Ser  
1115 1120 1125

Ser Met Phe Cys Leu Ser Pro Thr Asn Leu Leu Arg Arg Phe Cys  
1130 1135 1140

His Tyr Ile Val Thr Met Arg Tyr Phe Glu Val Val Ile Leu Val  
1145 1150 1155

Val Ile Ala Leu Ser Ser Ile Ala Leu Ala Ala Glu Asp Pro Val  
1160 1165 1170

Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys Tyr Leu Asp Tyr  
 1175 1180 1185

Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile Lys Met Ile  
 1190 1195 1200

Asp Leu Gly Leu Leu Leu His Pro Gly Ala Tyr Phe Arg Asp Leu  
 1205 1210 1215

Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val Ala  
 1220 1225 1230

Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys  
 1235 1240 1245

Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys  
 1250 1255 1260

Arg Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser  
 1265 1270 1275

Leu Lys Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met  
 1280 1285 1290

Phe Ile Phe Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe  
 1295 1300 1305

Phe Tyr Cys Thr Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg  
 1310 1315 1320

Gly Gln Tyr Leu Asp Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro  
 1325 1330 1335

Arg Gln Trp Lys Lys Tyr Asp Phe His Tyr Asp Asn Val Leu Trp  
 1340 1345 1350

Ala Leu Leu Thr Leu Phe Thr Val Ser Thr Gly Glu Gly Trp Pro  
 1355 1360 1365

Met Val Leu Lys His Ser Val Asp Ala Thr Tyr Glu Glu Gln Gly  
 1370 1375 1380

Pro Ser Pro Gly Tyr Arg Met Glu Leu Ser Ile Phe Tyr Val Val  
 1385 1390 1395

Tyr Phe Val Val Phe Pro Phe Phe Phe Val Asn Ile Phe Val Ala  
 1400 1405 1410

Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp Lys Val Met Ser  
 1415 1420 1425

Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile Asp Phe Ala  
 1430 1435 1440

Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn Arg Gln  
 1445 1450 1455

Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro Phe  
 1460 1465 1470

Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu  
 1475 1480 1485

Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu  
 1490 1495 1500

Lys Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys  
 1505 1510 1515

Val Leu Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp  
 1520 1525 1530

Ala Trp Asn Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr  
 1535 1540 1545

Asp Ile Leu Val Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn  
 1550 1555 1560

Leu Ser Phe Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu  
 1565 1570 1575

Leu Arg Gln Gly Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val  
 1580 1585 1590

Gln Ser Phe Lys Ala Leu Pro Tyr Val Cys Leu Leu Ile Ala Met  
 1595 1600 1605

Leu Phe Phe Ile Tyr Ala Ile Ile Gly Met Gln Val Phe Gly Asn  
 1610 1615 1620

Ile Ala Leu Asp Asp Asp Thr Ser Ile Asn Arg His Asn Asn Phe  
1625 1630 1635

Arg Thr Phe Leu Gln Ala Leu Met Leu Leu Phe Arg Ser Ala Thr  
1640 1645 1650

Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys Leu Ser Asn Gln  
1655 1660 1665

Ala Cys Asp Glu Gln Ala Asn Ala Thr Glu Cys Gly Ser Asp Phe  
1670 1675 1680

Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser Phe Leu  
1685 1690 1695

Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu Tyr  
1700 1705 1710

Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu  
1715 1720 1725

Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys Ala Trp  
1730 1735 1740

Phe Ala  
1745

<210> 5  
<211> 39  
<212> DNA  
<213> Homo sapiens

<400> 5  
tcttcctgtg ctcccttctc gcctggttcg catgaacat

39

<210> 6  
<211> 40  
<212> DNA  
<213> Homo sapiens

<400> 6  
gaatacgacc cggctgcgtg cgcctggttc gcatgaacat

40

<210> 7  
<211> 10  
<212> PRT  
<213> Homo sapiens

```

<400> 7
Leu Cys Ser Phe Leu Arg Leu Val Arg Met
1           5           10

<210> 8
<211> 10
<212> PRT
<213> Homo sapiens

<400> 8
Tyr Asp Pro Ala Ala Cys Ala Trp Phe Ala
1           5           10

<210> 9
<211> 28
<212> DNA
<213> Homo sapiens

<400> 9
gtttgggaat attgccctgg atgatgac          28

<210> 10
<211> 28
<212> DNA
<213> Homo sapiens

<400> 10
cttccccact gtcatctcat caggctta          28

<210> 11
<211> 26
<212> DNA
<213> Homo sapiens

<400> 11
atggtccgct tcggggacga gctggg            26

<210> 12
<211> 26
<212> DNA
<213> Homo sapiens

<400> 12
gcaccagtgg tcttggtcag ggtggt            26

<210> 13
<211> 26
<212> DNA
<213> Homo sapiens

<400> 13
tgcgaaccag gcgcacgcag ccgggt            26

```